SEQ ID NO.	BIALLELIC MARKER ID	ORIGINAL ALLELE	ALTERNATIVE ALLELE
1	20-828-311	C	T
1			·
1, 4	17-42-319	С	Т
1, 2, 4	17-41-250	С	Т
1	20-841-149	A	G
1	20-842-115	G	A
1	20-853-415	С	Т

Figure 1

SEQ ID no.	Biallelic Marker ID	Original Allele	Alternative Allele	Position Range of Preferred Sequences
1	20-828-311	C	T	739-1739
1	17-42-319	С	Т	10946-12958; 13470- 13526; 13641-13752
1	17-41-250	С	Т	14271-17969
1	20-841-149	A	G	41718-42718
1	20-842-115	G	A	44942-45942
1	20-853-415	С	Т	76558-77558
2	17-41-250	С	Т	1-1879
4	17-42-319	С	Т	1-1498; 1613-1724
4	17-41-250	C	Т	2243-3940; 3941-5381

Figure 2

A

SEQ ID NO.	POSITION OF CONFLICT	NUCLEOTIDE
1	13269 (SEQ ID No 1)	T (original)
4	1241 (SEQ ID No 4)	C (alternative)

B

SEQ ID NO.	POSITION OF CONFLICT	NUCLEOTIDE
1	13475 (SEQ ID No 1)	G (original)
4	1447 (SEQ ID No 4)	A (alternative)

Figures 3A, 3B

SEQ. ID. NO	POSITION RANGE O MICROSEQUENCING PRIMERS	COMPLEMENTARY POSITIN RANGE OF MICROSEQUENCING PRIMERS
1	1220-1238	1240-1258
1	12328-12346	12348-12366
1	15222-15240	15242-15260
1	42199-42217	42219-42237
1	45423-45441	45443-45461
1	77039-77057	77059-77077
4	300-318	320-338
4	3194-3212	3214-3232

Figure 4

SEQ. ID N0.	POSITION RANGE OF AMPLIFICATION PRIMERS	COMPLEMENTARY POSITION RANGE OF AMPLIFICATION PRIMERS
		1257 1277
11	929-949	1357-1377
1	12029-12050	12581-12603
1	14992-15012	15460-15482
1	42070-42090	42572-42591
1	45328-45347	45863-45883
1	76644-76664	77166-77185
4	1-11022	553-11575
4	899-11920	1441-12461
4	1246-12267	1632-12651
4	2964-13984	3432-14454

Figure 5

SEQ. ID NO	POSITION RANGE OF PFOBES
1	1227-1251
1	12335-12359
1	15229-15253
1	42206-42230
1	45430-45454
1	77046-77070
4	307-331
4	3201-3225

Figure 6

Alignment of ApoA IV-related cDNA with Human and Swine cDNA's

	AGACGTGAGCAGAGCAGATAATGGCAAGCATGGCTGCGGTGCTCACCTGGGCTCTGCGT_CTTCTTTCAGCGTTTTCAGCGT
ApoA IV related	CACCIC ACCIC ACCIC TOTAL CACCIC LOAT IS INCOME.
Human ApoA IV	AGTICCCACTGLAGGG-AGCTG-CCTGAGAAACCTCTCCTCCACGGAGG 39
Swine ApoA IV	
ApoA IV related	CACCCAGGCACGGAAAGGCTTCTGGGACTACTTCAGCGAGCCGGGACCAGAGCGGGGAGCAGGTGGAGCAGATCGATC
Human ApoA IV	CATCCAG-TGTGGGAAGAAACTCCTCCAGCCCAGCAAGCAGCT-CAGGATGTTCCTGA
Swine ApoA IV	•
ApoA IV related	AGAAGATGGCTGGGGAGCCCGGACCCTGAAAGACAGCCTTGAGGAAGACCCTCAACAATATGAACAAGTTCCTGGAAAAAC 239
Human ApoA IV	
Swine ApoA IV	AGGCTGTGGCCTGGCCCTGGTGGCCGTCACCGGTGCCCGGGCTGAGGTCAATGCCGACCAAG 162
•	CTGAGGCQTCTGAGTGGGAGGAGGAGTCCTCGGCTCGCACAGACCCGGTGGGCATGCGGGGGAGCAGCAGGAGGAGTT 319
ApoA IV related	TOTAL GOOGLACE TOTAL GOOGLACE TOTAL GOOGLACE TO TOTAL GOOGLACE TO TOTAL GOOGLACE TO THE
Human ApoA IV	TGGCTACHG-TGAT-GTGGGACTACTTCAGCCAG-CTGGGCAGCAATGCCAAGAAGGCTGT-GGAACATCT 229
Swine ApoA IV	
ApoA IV related	GGAGGAGGTGAAGGCTCGCGTCCAGCCCTACATGGCAGAGGGGCAGGAGGGGCTGGGATTTTGGAGGCTTTGCGGC 399
Human ApoA IV	CCAG-AAATCTGAACTCACCCAGCAACTCAATGCCCTCTTCCAGGACAAACTTGGAG316 CCAG-AAGTCTGAGCTCACCCAGCAGCTCAACACTCTCTTCCAGGACAAACTTGGGG285
Swine ApoA IV	
ApoA IV related	AGCAACTGAAGCCCTACACGATGGATCTGATGGAGCAGCTGCCCCTTGCCGGTGCAGGAGCTGCAGGAGCAGTTGCGCGTG 479
Human ApoA IV	TOTAL STATE OF CONTROL CONTROL CONTROL CONTROL OF THE CONTROL OF T
Swine ApoA IV	AAGTGAACACCTACACGGAGGACCTGCAGAAGAAGCTGGTGCCCTTTGCCACGGAGCTGCATGAACGCCTG 356
•	GTGGGGGAAGAAGAAGACCAAGIITGCTGGGGGGGGGGGACGAAGAGCCTTGGGGGGGAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGA
ApoA IV related	CARCOLOGICACA - TACTO - AACCAG GAGATTGGGAAGGAGCTGAGGGCCCGGC 452
Human ApoA IV	ACCAAGGACTGAGAGAAGCTGAAGGAGGAGATTGGAAGGGAGCTGGAGGAGCTGCGAGCCCGGCT 421
Swine ApoA IV	
ApoA IV related	GGTGCACCACACCCCCATACCCCCATACCCCATACCCCATACCCCATACCCCATACCCCATACCCCACACCCCCC
Human ApoA IV	GCTGCCCCACGCTACCGAGGTGAGCCAGAAAATCGGAGACAACGTGCGGGAGCTGCAGCAG-CGCCT- 487
Swine ApoA IV	
ApoA IV related	CAGGAGTTGGAACCGAGTTGTCGCTCCGGAAGGCCCCCGGCCAGGGCCCCCCAGTTGCTGCCAGGTGCTGTGTGGCCCCAGCTGCTGCAGGTGCTGCTGCCCCCAGCTGCAGCTGCTGCAGGTGCTGCGCCCAGCTGCAGCTGCGCCCAGCTGCAGCTGCGCCCAGCTGCAGCTGCGCCCAGCTGCAGCTGCGCCCAGCTGCAGCTGCGCCCAGCTGCAGCTAGCAGCTGCGCCCAGCTGCAGCTAGCAGCTGCGCCCAGCTGCAGCTAGCAGCTGCGCCCAGCTGCAGCTAGCAGCTGCGCCCAGCTGCAGCTAGCAGCTGCGCCCAGCTGCAGCTAGCAGCTGCGCCCAGCTGCAGCTAGCAGCTGCGCCCAGCTGCAGCTGCAGCTAGCAGCTGCGCCCCAGCTGCAGCTAGCAGCTGCGCCCCAGCTGCAGCTAGCAGCTGCAGCTAGCAGCTGCAGCTAGCAGCAGCTAGCAGCTAGCAGCTAGCAGCTAGCAGCAGCAGCAGCAGCTAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCA
Human ApoA IV	GGAGCC-CTHACGCGACCAGCTGCGCACCCAGGTCAACACG-CAGGCC-GAGCAGCTGCGGCGC-CAGCTG 585GGGGCC-CTTTACGGGAGGGCTGCGCACCCAGGTCAACACC-CAGGTT-CAGCAGCTGGAGCGC-CAGCTG 554
Swine ApoA IV	
ApoA IV related	GAAGOTTAACGCTCAAGGCCAAGGCCTGCACGCACGCAGAGCCAGAGCTGCGCCGAAGAGCTCAGCAGAGCTGAGCAGAGCAGAGCTGAGCAGAGCTGAGCAGAGCAGAGCTGAGCAGAGCAGAGCTGAGCAGAGAGCAGAGAGCAGAGAGCAGAGAGCAGAGAGCAGAGAGCAGAGAGCAGAGAGCAGAGAGCAGAGAGCAGAGAGCAGAGAGAGAGCAGAGAGAGAGCAGAGAGAGAGAGCAGAGAGAGAGCAG
Human ApoA IV	GONDOCA CARDOCA CARDOCA CARDA CONTROL TICOCOCAGA A C
Swine ApoA IV	ACCCCTACGCALAGCGCATGGAGTCCGTGC-TACGGCAGAACATCCGCAACCTGGAGGCCTCCGTGGCA 623
•	cqrii-redalqqacriqqqacriqqqaaqqaqqqqqqqqqqqqqqqqqqq
ApoA IV related	CACCACACCACCACCACACCACACCACACCACCACCACC
Human ApoA IV	CCCTATGGGGATGAGTTCAAGGCCAAGATCGATCAGAATGTGGAGGAGCTCAAGGGGA 681
Swine ApoA IV	
ApoA IV related	GGGTTTGCGGCAGGACACCTAGCTGGAGATAGCTGGCTTCAGTGGGCCACCACCACCACCACCACCACCACCACCACCACCAC
Human ApoA IV	-GCCTTACGCCCTATGCGGAGGAGCTCAAGGCAAGATCGATCAGAACGTGGAGGAGCTGCGGC-GC 746
Swine ApoA IV	
ApoA IV related	AGTTGGGGCGACCTCCAGCAGGGCACAGTGCTTTGCCCCCAGAGTTTTCAACAAAACAGACAG
Human ApoA IV	AGCTTG-GCTCCCTATGCTCAGGACACGCAGGAGAAGCTCAACCACCAGCTTGAGGGCCTGA838 AGCCTG-GCCCCCTATGCGCAGGACCTCCAGGAGAAGCTCAACCACCAGCTCGAGGGCCTGG807
Swine ApoA IV	
ApoA IV related	CTGCAGGCCGTTTGGGATGACCTGTGGGAAGAATDACTCACAGGCTTCATGACCAGGGCGAGAGCCATCTGGGGGACCC 1117
Human ApoA IV	ACA
Swine ApoA IV	CCTTCCAGATGAAGAAGCAGGAGAGGAGCTGAAGGCCAAGATCTCGGCCAA-TG
	THEAGGATCHAGCTGCCCAGGCCGATTCCGAGCTCCTTGTCTGGGGAGCCTTGGGTTGAGCGTCTAGCATGGTTCAGTG 1197
ApoA IV related	CCCACA
Human ApoA IV	CCGAGGAGCTGCGCAGAAGCTGGTGCCGGTGGCCGAGAACGTGCATGGC-CAT 914
Swine ApoA IV	
ApoA IV related	CITCHANGTGGGCTGTTGGGTGGAGGTGGAAGGTCCTGTGCAGGAGAGGGGAGGCCACCAAAGGGGGTGCTGTCTCCTGC 1277 CTGAGGGGCAACACCGAG-GGGCTGCAGAAGTGACTG-GCAGAGCTGGGTGGGCACCTGGACCAGCAGGTGGAGG 1018
Human ApoA IV	CTGAGGGGCAACACCGAG-GGGCTGCAGAAGTGACTG-GCAGAGCTGGGTGGGCACCTGGACCAGCAGGTGGAGG CTGAAGGGCAACACCGAG-GGGCTGCAGAAGTGCTG-CTGGAGCTGAGAAGCCACCTGGACCAGCAGGTGGAGG-987
Swine ApoA IV	
ApoA IV related	ATATCCAGGCTGCTGCGACTCCCCGATGTGGATGCATTACATTCACCAGGCTTTGCAAACCCAGGCTCCCAGTGCTCATT 1357
Human ApoA IV	- CONTROL OCCOUNTS CARCULAR - COCCENTRAL - ACAAAGCCCTG GTGCAGC ACAAAGCCCTG
Swine ApoA IV	AGTICCOCCITAAGGTGGAGCCCTA - CGGGAGACCTTCA ACAAAGCTTTG GTGCAGC AG G 1047

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Alignment of ApoA IV-related cDNA with Human and Swine cDNA's

ApoA IV related	TEGGRATICTCATGAGTTACTCCATTGAAGGGTGAGGGAGTAGGGAGGAGAGGCACCATGCATG	1437
Human ApoA IV	TCCA BCACCTCAGGACGAAACTGGGCCCCCATGCGGGGGACGTGGAAGGCCACTTG	1134
Swine ApoA IV	TGGAGGATCTCAGGGAGAAGCTGGGCCCTTTGGCGGGGGACGTGGAGGGCCACCTG-	1103
Swille ApoA IV		
ApoA IV related	NACOCICTIFIC COTTO A TOCK OF THE CONTROL OF THE CON	1517
Human ApoA IV	DOCTTCCTCCAGAAGGACCTGAGGGACAAGGTCAACTCCTTCTTCAG-CACCTTCAAGGAGA	
•	-AGCTTCCTGGAGAAGGATCTGAGGGACAAGGTCAACACCTTCTTCAG-CACCCTCAAAGAGG	1164
Swine ApoA IV		
ApoA IV related	CCACTGCTACAGCTGCTCCACAGAGAGGACGACTTGTCTCCCCAGGGCTGCCATGGCAGCTATCAGGGGAATAGAAGGGA	1597
Human ApoA IV	AAGAGAGCCA-GGACAAGACTT-CTCTCCCTCCCTGAGCTGGAGCAACAGCAGGAACAGCATCAG	
Swine ApoA IV	GCdGAGCCA-GGddCAGAGC-CACGCCCTCCCTGCA	1199
Swille Apon IV		
ApoA IV related	MANACACINATICATICATICATICATICATICATICATICATICATIC	1677
Human ApoA IV	WACCACONG ACTION ACCOUNTS ACCO	1335
Swine ApoA IV	CAGGAGAAGGCCAGGCCCCTTTGGAGGGCTGAGCTGCCCCTGGTGCTCCCACCCCAC	1262
Swine ApoA iv		1757
ApoA IV related	TIGGGTTGTGATAGGAGAGGGCAGAGCCCATGTTTTCCTGAGATAGGTTTACACCTAAATAAGGGACTGAACCCCAACT	1376
Human ApoA IV	TGGACGTCTGT-CTGTCCCAA-	1304
Swine ApoA IV	ACCTGCCCTGCCCT-GCCCCCTGTCTGTCTGT-CTGTCCCAA	1304
Dwille Lipoliti		1027
ApoA IV related	GTGGGAGGTGGTTAAAGCCTGTGGGAAGCATACTGTGTGCTCAGGCGGTTCCCAAGCTGCTCTGGGTTCCCAAGTTTG	103/
Human ApoA IV	AGAAGTTC-TGGTATGAACTTGAGGACACATGTCCAGTGGGAGGTGAGACCCACCTCTGAATATTCAATA	1372
Swine ApoA IV	AGCAGTTC-TTGTACHAAGCHAGGGATACATGTCCAGTGGACCGTGACACTACCTCTCCATACTCAATA	13,2
Swille Pipori IV		1879
ApoA IV related	AAGCTAGACTTCTGGCTCAAATGAATAGATGTTTATGATA	1466
Human ApoA IV	AAGCTGCTGAGAATCTAGCCEC	1388
Swine ApoA IV	AAGCTGCTGAGAAACT	

Alignment of ApoA IV-related protein with Human and Swine ApoA IV

ApoA IV related	MASMAAVLTWALALLSAFSATGARKGFWDYFSGTISGD-KGRVEQIHQQKMAREPATI-KHSILEQDLNNMNKFLEKL 74
Human ApoA IV	MFLKAVVLTLALVAVAGARAEVSADQVATVMWDYFSQLSNNAKEAVEHLQKSELTQQLNALFQDKLGEVNTYAGDLQKKL 80
Swine ApoA IV	MFLKAVVISLALVAVTGARAEVNADQVATVMWDYFSQLGSNAKKAVEHLQKSELTQQLNTLFQDKLGEVNTYTBDLQKKL 80
ApoA IV related	RHLSGSBAPRIIPQDPVGMRRQLQBELEBVKARIIQFYMABAHELVGWNIBGIIRGQIKPYJMDIMEGVALRVGBLGBGIRVV 154
Human ApoA IV	VPFATELHERIAKDSEKLKEEJGKELEELRARLLPHANEVSQKIGDNIRBLQQRIBPYADQLRTQVNTQABQIRRQUTPY 160
Swine ApoA IV	VPFATELHERITKDSEKLKEEJRRBLEELRARLLPHATEVSQKIGDNVRELQQRIGFFIGGLRTQVNTQVQQLQRQIKPY 160
ApoA IV related	GBDTKAQHLGGVDEAWALHQGHQSRVVHHTGRFKBELFHPYABSHVSGHGRHVQBHHRSVAFHAPASPARHSRCVQV 230
Human ApoA IV	AQRMBRVLRBNADSHQASHRBHADELKAKIDQNVEELKGRLTPYADHFKVKIDQTVEELRRSLAPYAQHQEKLNHQLEG 240
Swine ApoA IV	AERMBSVLRQNIRNUBASVARYADHFKAKIDQNVEELKGSLTPYABELKAKIDQNVEELRRSLAPYAQDVQEKLNHQLEG 240
ApoA IV related	USRKLTLKAKATHAR 100N-100REBUSRAFAGTGTEBGAGPDPOMUSEBVRGRLOAFRODTVLOIAAFTRAIDG 305
Human ApoA IV	LTFOMKRNAEBLKARISASABELRGRLAAFLABDVRGNURGNTEGLOKSUAEUGGHLDQOVEBFRRKVEPYGBNFNKALVG 320
Swine ApoA IV	ЦАFOMKKQAEBLKAKISANADELRGKUVBVABNVHGHUKGNTEGLOKSULBURSHLDQOVEBFRLKVEPYGBTFNKALVG 320
ApoA IV related	ETBEVOGOLARPPPGHSAFAPEFQQTDSGKVLSKLQARLDDLWEDIRHSIHDGGHSHLGDR 366
Human ApoA IV	QWBQLRTKLGPHAGDVEGHLSFLEKDLRDKVNSFFSTFKBKOCKILSLFELBQQQBQHQBQQQBQVQMLAPLES 396
Swine ApoA IV	QVBDLRQKLGPLAGDVEGHLSFLEKDLRDKVNTFFSTLKBBASQGQSQALFAQBKAQAPLBG 382

Figure 8

Alignment of ApoA IV-related cDNA with Rat RAP3 cDNAs

ApoA IV related	AGACGTGAGCAGAGAGATAATGGCAAGCATGGCTGCCGTGCTCACCTGGGCTCTGGCTCTTCTTTCAGCGTTTTCGGCC	B0
Rat RAP3 A	GCATCGTGGAAAGCATGGCTGCCGTCATCACCTGGGCACTCGCCCTCCTCTCAGTGTTTGCAACT	65
Rat RAP3 B	GCATCGTGGAAAGCATGGCTGCCGTCATCACCTGGGCACTCGCCCTCCTCAGIGITIGCAACI	65
ApoA IV related	ACCCAGGGACGGAAAGGCTTCTGGGACTACTTQAGCCAGACCAGCGGGGACAAAGGCAGGGTGGAGCAGATCCATCAGCA	160
Rat RAP3 A	CTACACCCGACGAAGAGCTTCTCGGAGTACTTCGGCCAGAACAGCCAGGGCAAAGGCATGATGGGCCAGCAGCA	139
Rat RAP3 B	GTACAGGCGAGGAAGAGCTTCTGGGAGTACTTCGGCCAGAACAGCCAGGGCAAAGGCATGATGGGCCAGCAGCA	139
ApoA IV related	GAAGATGGCTTGGCGAGCCCGCGACCTTGAAAGACAGCCTTGAGCAAGACCTCAACAATATGAACAAGTTCCTGGAAAAAGC	240
Rat RAP3 A	GAAGATGGCTGGGAAGGAGAGCCTGAAAGGTAGCTTGGAGCAAGACCTCTACAATATGAACAATTTCCTAGAAAAGC	213
Rat RAP3 B	GAAGCTGGCACAGGAGAGCCTGAAAGGTAGCTTGGAGCAAGACCTCTACAATATGAACAATTTCCTAGAAAAAGC	213
ApoA IV related	ITGAGGCCTCTGAGTGGGAGCGAGCCTCCTCGGCTCCCACAGGACCCGGCTGGGCATGCGGCAGCTGCAGGAG	314
Rat RAP3 A	TGGGACCCTTGAGAGAGCCTGGGAAGGAGCCTCCTCGGCTGGCACAGGATCCAGAAGGCATTCGGAAGCAGTTGCAGCAA	293
Rat RAP3 B	TGGGACCCTTGAGAGAGCCTGGGAAGGAGCCTCCTCGGCTGGCACAGGATCCAGAAGGCATTCGGAAGCAGTTGCAGCAA	293
ApoA IV related	GACTITGGAGGACGTGAAGGTTCGCCTCCAGCCCTACATGGCAGAGGCGCAGCAGGTGGTGGGCTGGAATTTTGGAGGGCTT	394
Rat RAP3 A	CACCTCCACGA ACTGAGCA CACGCCTGGAGCCCTACATGGCTGCAAAGCACCAGGAGGTCGGCTGGAACCTGGAGGGCT	3/3
Rat RAP3 B	GAGCTGGAGGAAGTGAGCACACGCCTGGAGCCCTACATGGCTGCAAAGCACCAGCAGGTCGGCTGGAACCTTGGAGGGCTT	373
ApoA IV related	GOGGCAGCAACTGAAGCCCTACACGATGGATCTGATGGAGCAGGTGGCCCTGCGCGTGCAGGAGCAGCAGTTGG	474
Rat RAP3 A	CACCACCACTAGA AACCCTACACGTCGAGCTGATGGAGCAGGTAGGCCTGAGCGTGCAGGATCTGCAAGAACAGCTGU	453
Rat RAP3 B	GAGGCAGCAGTTGAAACCCTACACGGTCGAGCTGATGGAGCAGGTAGGCCTGAGCGTGCAGGATCTGCAAGAACAGCTGC	453
ApoA IV related	GGGTGGTGGGGAAGACACCAAGGCCCAGTTGCTGGGGGGCGTGGAGGAGGCTTGGGCTTTGCTGCAGGGACTGCAGAGC	554
Rat RAP3 A	CATCCTCCGA & ACCCACCAGGCCCAGCTCCTGGGGGGGGGTGAGGCGATGAGCCTGCTGCAGGATATGCAAAGT	533
Rat RAP3 B	GCATGGTGGGAAAAGGCACCAAGGCCCAGCTCCTGGGGGGGCGTGGATGAGGCGTGAGCCTGCTGCAGGATATGCAAAAGI	533
ApoA IV related	COCTCGTGCACCACACCCCTTCAAAGACCTCTTCCACCCATACGCGGAGAGCCTGGGGGAGAGCGGGATCGGGGCCCA	634
Rat RAP3 A	CCACTCCTCCACCATACCGACCGAGTCAAAGAACTCTTCCACCCTTATGCAGAACGCTTGGTGACTGGAATTGGGCACCA	913
Rat RAP3 B	CGAGTGCTGCACCATACGGACCGAGTCAAAGAACTCTTCCACCCTTATGCAGAACGCTTGGTGACTGGAATTGGGCACCA	613
ApoA IV related	GETGCAGGAGCTGCACCGGAGTGTGGCTCCGCACGGCCCAGCCCCGCGGGCTCAGTCGCTGCGTGCAGGTGCTGT	714
Rat RAP3 A	FOTCO CONCETCO CCCO CTCTTCCTCTCACCCACTTGCCACCCCCGCGAGACTCACTC	693
Rat RAP3 B	TGTGCAGGAGCTGCACCGGAGTGTTGCTCCTCACGCAGTTGCCAGCCCCGCGAGACTCAGTCGCTGCGTGCAGACCCTGT	093
ApoA IV related	CCCCGAAGCTCACGCTCAAGGCAAGACCCCTGCACGCACCACCAGCAAACCTGGACCAGCTGCCGGAAGAGCTCAGC	794
Rat RAP3 A	CCCACAAACTCACACGTAAGGCGAAGGACTTGCACACCAGCATCCAACGCAACCTGGATCAGCTGCGAGATGAGCTCAGT	7/3
Rat RAP3 B	CCCACAAACTCACACGTAAGGCGAAGGACTTGCACACCAGCATCCAACGCAACCTGGATCAGCTGCGAGATGAGCTCAGT	//3
ApoA IV related	AGAGCGTTTTGCAGGCAdTGGGACTGAGGAAGGGGCGGGCCCGGGACCCCCAGATGCTCTCCGAGGAGGTGCGCCA	868
Rat RAP3 A	A COTTCATCOCTET CAGCA CAGA CEGGGGAGACAACAGAGACTCCCTGGACCCTCAAGCTCTCTCT	853
Rat RAP3 B	ACCTTCATCCGTGTCAGCACAGACGGGGCAGACAACAGAGACTCCCTGGACCCTCAAGCTCTCTGACGAGGTCCGCCA	853
ApoA IV related	GGGACTTCAGGCTTTCCGCCAGGACACCTACCTGCAGATAGCTGCCTTCACTCGCGCCCATGGACCAGGAGAGTGAGGAGG	948
Rat RAP3 A	CACACTCCACGCTTTTCGACATGACACCTGCCTGCAGATCGCTGCATTCACTCAGGCCATTGACCAGGAGACCGAGGAAA	933
Rat RAP3 B	GAGACTCCAGGCTTTTCGACATGACACCTACCTGCAGATCGCTGCATTCACTCAGGCCATTGACCAGGAGACCGAGGAAA	933
ApoA IV related	TCCAGCAGCAGCTGGCGCCACCTTCCACCACACTGCCACACTTCCCCCCAGAGTTTCAACAAACA	1028
Rat RAP3 A	TRECACENCE CONCERCE CONCERCE ACCOUNT A	1012
Rat RAP3 B	TCCAGCACCAGCTGGCACCACCCCCGGCTAGCCACAGCGCCTTCGCTCCAGAGTTGGGACACTCAGACAGTAATAAGGCC	1013
ApoA IV related	CTGAGCAAGCTGCAGGCCCATCTGGATGACCTGTGGGAAGACATCACTCAC	1108
Rat RAP3 A	CTGAGCAGACTGCAGAGCCGGCTGGACGACCTCTGGGAAGATATTGCCTATGGCCTTCATGACCAGGGCCATAGTCA	1090
Rat RAP3 B	CTGAGCAGACTGCAGAGCCGGCTGGACGACCTCTGGGAAGATATTGCCTATGGCCTTCATGACCAGGGCCATAGTCA	1090
ApoA IV related	GGGGGACCCTGAGGATCTACCTGCCCAGGCCCCTTTCC-CAGCTCCTTGTCTGGGGAGCCTTGGCTCTGAGCCTCTAGCA	1187
Rat RAP3 A	CANTA ACCOMPAGGETC ACT CAGGTTA ACT CTGCAGCT CGTTGT CTGGA CCCTGAGCCT CAGCA	1122
Rat RAP3 B	-GAATAACCCTGAGGGTCACTCAGGTTAACTCTGCAGCTCGTTGTCTGGACCCTGAGCCTTCAGCA	1133

Figure 9 A

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Alignment of ApoA IV-related cDNA with Rat RAP3 cDNAs

ApoA IV related	TEGTTCAGTCCTTGAAAGTGGCCTGTTGGGGGGGGGGGGG	1266
Rat RAP3 A	TGGCCTAATAGGCAGAGGGTGGAGGGTCCTGCATACTATTGGCGAGGCCACCAAAGGTGCTG	1217
Rat RAP3 B	TGGCCTAATAGGCAGAGGGTGGAGGGTCCTGCATACTATTGGCGAGGCCACCAAAGGTGCTG	1217
	The state of the s	1246
ApoA IV related	CTGTCTCGTGCATATTGCAGCCTCCTGGGACTCCCGAATCCTGGATGCATTACATTCACCAGGGTTTGCAAAGCAGGGTCCCTG-CCCCAACCTGTCGCCAAACCCCCACTCAGGTGCATTACACTCAGTAGGTTTGCCAAAGACAGGTTCG	1295
Rat RAP3 A	CTG-CCCCAACCTGTCTGGCCTCCT-CAACTCCCCACTCAGGTGCATTACACTCAGTAGGTTTGGCAAA	1285
Rat RAP3 B	CTG-CCCCAACCTGTCTGGCCTCCT-CAACTCCCCCACTCAGGTGCATTACACTCAGTAGGTTTGGCAAA	
ApoA IV related	CAGTGCTCATTTGGGAATGCTCATGAGTTACTCCATTCAAGCGTGAGGGAGTTAGGGGAGGGA	1426
Rat RAP3 A	GCTGCTCATTTGGGA-TCCTAAGGAGCAAGAGTG-GCGTGAAGGGAGTGGGGAG-ATGGTGTGGGGGG	1361
Rat RAP3 B		1285
Kat KAF3 B		
ApoA IV related	TGATTATCTGCAAGCC - TGTTTGCGGTGATGCTGGAAGCCTGTGCCACTACATCCTGGAGTTTGGCTCTAGTCACTTGT	1504
Rat RAP3 A	AGACTGACTGCAAGCCAGTIACTTGAC COTTGCTAGAAACCTGTGTCACTACAACCTGGAGCCCGGCTCCTATTACTTGA	1440
Rat RAP3 B		1285
ApoA IV related	GGGTGCCTGGTGGGACTGCTACAGCTGGTQACAGAGAGGAGGACTTGTCTCCCCAGGGCTGCCATGGCAGGTATCAGG	1584
Rat RAP3 A	TGCCTGATGGTGGCTGTTATAGTCGGTGTACAGAGGGGAAGTCCTGTCTCCCCAGGGTTGTCATGACAGGCTGTGTT	1285
Rat RAP3 B		1203
ApoA IV related	GGAATAGAAGGGAGAAAGAGAATATCATGGGGAGAACATGTGATGTGTGTAATATCCCTGCTGCTCTGATG	1658
Rat RAP3 A	GGAAGAGAGCAGGAGAACATGACGTATGATGGAGTGTGTACATCCCTGCCAGTGGTCCTGCTGCGGGGAATCAGTGATG	1597
Rat RAP3 B	bondano de de la companya de la comp	1285
Kai KAP3 D		
ApoA IV related	CTGGTGGGTACGAAAGGTGTGGGCTGTGATAGGAGAGGGCAGAGCCCATGTTTCCTGACATA	1720
Rat RAP3 A	GGATAAATGTGTGGATCCCTGGAGTGGTCCTGGTGGGGGATGAGTGATGGGATGGGGCAGAGCCGGTATTTCCTTAGAGA	1677
Rat RAP3 B		1285
		1799
ApoA IV related	GCTCTAGACCTAAATAAGGGACTGAACCCTGCCAACTGTGGGAGCTCCTTAAA-CCCTCTGGGGAGCATACTGTGTGGTG	1754
Rat RAP3 A	ACTCTA-ACCCAAATAAGGAACTGAGCCCTGT-GGAGTGAGGCCTTCTGAAAACCCTGTACATAGCAAACTGTGTGGCG	1285
Rat RAP3 B		1203
ApoA IV related	TOCCCATC-TCCAGCCCTCCCTCTCCGTTCCGAAGTTCAAGCCTAGACTTCTCGCTCAAATGAAATAGATCTTTATGAT	1878
Rat RAP3 A	TOTTCATCATGCAGTCCCCCACCTCCTGATTCTCCCGGATCGAACT GACTTTTCGTTTCGAAATGAAATAGACGCTCATGAT	1832
Rat RAP3 B		1285
ApoA IV related	A	1879
Rat RAP3 A	GGAAAAAAAAAAA	1850 1300
Rat RAP3 B	AAAAAAAAAAAA	1300

Alignment of ApoA IV-related protein to Rat RAP3 proteins

78	158	238	315	366
73	153	233	312	367
73	153	233	312	367
ed mas maavijtwalalisafisatoarkofwdyfisotsotsokomkols	GSBAPRLPODPYGMRROLOEBLEEVKARLOPYMABAHELVGWNLEGLRQOLKPYTMDLMEGVALIRVOELOEOLRYVGEDF	ed kaqılıggvdeamalıqgılosrvvhhttgrekelfhpyaeişlvisgigihvqelhrsvaphalarsparisrcvqvlsirkitik	ed ARALHARITQONLDQLREELSRAFAGIGIEEGAGPDPOMLSEEVRQRLQAFRQDTYLQIAAFTRAIDOETEEVOOLA	ed ppp pighsafappel <u>pootdsigkvlsklodarlddenthal</u> hdoghsundens <u>hlgdp</u>
maavitwalalisyfatvoarksfweyfgonsogkommgoooklaoeslkgsleodlynmnnflekloplrep	GKEPPRLAQDPEGIRKOLQQELEEVSTRLEPYMAAKHQQVGWNLEGLRQQLKPYTVELMEQVGLSVQDLQEQLRWVGKGT	kaqılggvdeamsılqdmqsrvlhhtdrvkelfhpyaerlvtgighhvqelhrsvaphavasparisrcvqtlshkltrk	AKDLHTSIQRNLDQLRDELS-TFIRVSTDGADNRDSLDPQALSDEVRQRLQAFRHDTYLQIAAFTQAIDQETEEIQHQLA	ppppshsafappelghsdsnkalsrlosrlddiwediayglhdoghsonndeghsg
maavitwalalisyfatvoarksfweyfgonsogkommgoooklaoeslkgsleodlynmnnflekloplrep	GKEPPRLAQDPEGIRKQLQQELEEVSTRLEPYMAAKHQQVGWNLEGLRQQLKPYTVELMEQVGLSVQDLQEQLRMYGKGT	kaqılggvdeamsılqdmqsrvlhhtdrvkelfhpyaerlvtgighhvqelhrsvaphavasparlsrcvqtlshkltrk	AKDLHTSIQRNLDQLRDELS-TEIRVSTDGADNRDSLDPQALSDEVRQRLQARRHDTYLQIAAFTQAIDQETEEIQHQLA	pppshsafapelghsdsnkalsrlosrlddiwediayglhdoghsonndeghsg
ApoA IV related	ApoA IV related	ApoA IV related	ApoA IV related	ApoA IV related
Rat RAP3 A	Rat RAP3 A	Rat RAP3 A	Rat RAP3 A	Rat RAP3 A
Rat RAP3 B	Rat RAP3 B	Rat RAP3 B	Rat RAP3 B	Rat RAP3 B

Figure 10

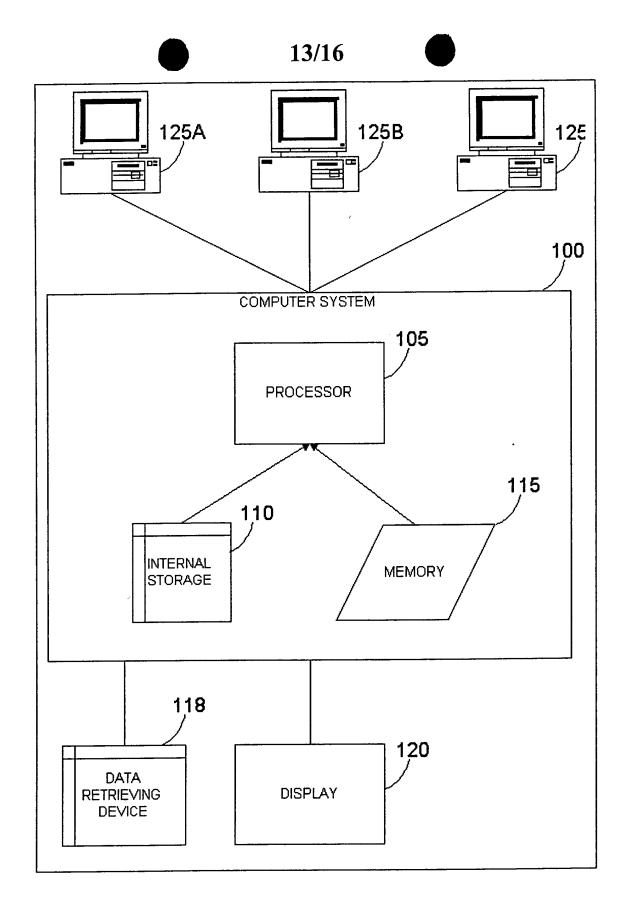


Figure 11

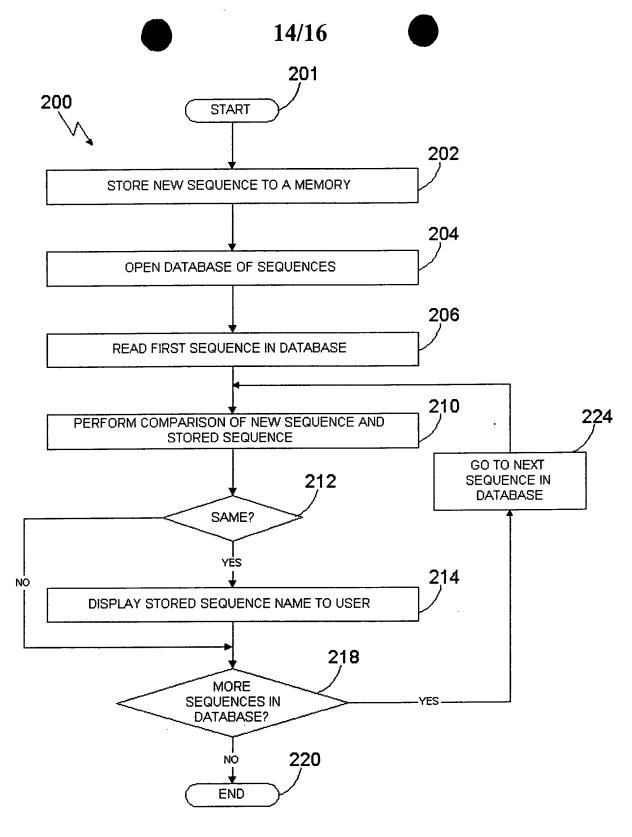


Figure 12

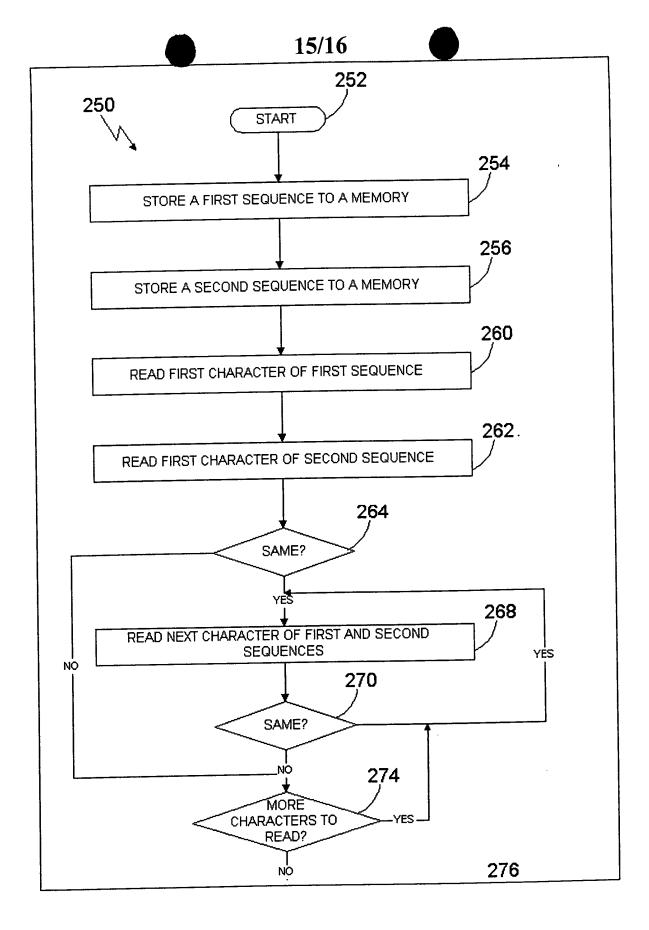


Figure 13

